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 Lys Ile Ser Asn His Gly Ser Leu Arg Val Ala Lys Val Ala Tyr Pro
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477

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tct Ser 115	Ата	tat Tyr	gct Ala	gat Asp	gat Asp 120	gtt Val	gaa Glu	aag Lys	Ser	gct Ala 125	Gln	gga Gly	ato	cgg	gaa Glu 130	621
Leu	Leu	Asp	Val	Ala 135	Lys	Gln	Asp	Ile	Pro	Phe	Asp	Phe	Trp	Lys 145		669
inr	PIO		150	Leu	Lys	Ala	Thr	Ala 155	Gly	Leu	Arg	Leu	Leu 160	Pro	Gly	717
GIU	гув	gcc Ala 165	GIN	ьуs	Leu	Leu	Gln 170	Lys	Val	Lys	Glu	Val 175	Phe	Lys	Ala	765
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gcc	ser	245	Pro	GIY	Tyr	Leu	250	Ala	Leu	Arg	Met	Phe 255	Asn	Arg	Thr	1005
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arg 275	ьеu	мта	тте	Leu	280	GIY	Val	Glu	Gly	Gln 285	Pro	Ala	Lys	Asp	Gly 290	1101
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Thr Ala Ala Asp Gly His Glu Val Phe Tyr Gly Ile Met Phe Asp Ala 65 70 75 80

Gly Ser Thr Gly Thr Arg Val His Val Phe Gln Phe Thr Arg Pro Pro 85 90 95

Arg Glu Thr Pro Thr Leu Thr His Glu Thr Phe Lys Ala Val Lys Pro 100 105 110

Gly Leu Ser Ala Tyr Ala Asp Asp Val Glu Lys Ser Ala Gln Gly Ile 115 120 125

Arg Glu Leu Leu Asp Val Ala Lys Gln Asp Ile Pro Phe Asp Phe Trp $130 \\ 135 \\ 140 \\ $

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- Leu Gln Ala Ser Pro Pro Gly Tyr Leu Thr Ala Leu Arg Met Phe Asn 245 250 255
- Arg Thr Tyr Lys Leu Tyr Ser Tyr Ser Tyr Leu Gly Leu Met $260 \hspace{1cm} 265 \hspace{1cm} 265 \hspace{1cm} 270 \hspace{1cm}$
- Ser Ala Arg Leu Ala Ile Leu Gly Gly Val Glu Gly Gln Pro Ala Lys 275 280 285
- Asp Gly Lys Glu Leu Val Ser Pro Cys Leu Ser Pro Ser Phe Lys Gly 290 300
- Glu Trp Glu His Ala Glu Val Thr Tyr Arg Val Ser Gly Gln Lys Ala 305 310 315
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Leu Val Val Leu Leu Val Ser Ile Val Val Leu Val Ser Ile Thr Val
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Ser Ile Val Val Leu Val Ser Ile Thr Val Ile Gln Ile His Lys Gln 35 40 45

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Ser Ser Arg Thr Thr Val Tyr Val Tyr Gln Trp Pro Ala Glu Lys Glu 65 70 75 80

Asn Asn Thr Gly Val Val Ser Gln Thr Phe Lys Cys Ser Val Lys Gly 85 90 95

Ser Gly Ile Ser Ser Tyr Gly Asn Asn Pro Gln Asp Val Pro Arg Ala 100 105 110

Phe Glu Glu Cys Met Gln Lys Val Lys Gly Gln Val Pro Ser His Leu 115 120 125

His Gly Ser Thr Pro IIe His Leu Gly Ala Thr Ala Gly Met Arg Leu 130 135, 140

Leu Arg Leu Gln Asn Glu Thr Ala Ala Asn Glu Val Leu Glu Ser Ile 145 150 155 160

Gln Ser Tyr Phe Lys Ser Gln Pro Phe Asp Phe Arg Gly Ala Gln Ile 165 170 175

Ile Ser Gly Gln Glu Glu Gly Val Tyr Gly Trp Ile Thr Ala Asn Tyr 180 185 190

- Leu Met Gly Asn Phe Leu Glu Lys Asn Leu Trp His Met Trp Val His
- Pro His Gly Val Glu Thr Thr Gly Ala Leu Asp Leu Gly Gly Ala Ser 210 215 220
- Thr Gln Ile Ser Phe Val Ala Gly Glu Lys Met Asp Leu Asn Thr Ser 225 230235235
- His Ser Phe Gln Cys Tyr Gly Arg Asn Glu Ala Glu Lys Lys Phe Leu 260 265 270
- Ala Met Leu Leu Gln Asn Ser Pro Thr Lys Asn His Leu Thr Asn Pro 275 280 285
- Cys Tyr Pro Arg Asp Tyr Ser Ile Ser Phe Thr Met Gly His Val Phe 290 295 300
- Asp Ser Leu Cys Thr Val Asp Gln Arg Pro Glu Ser Tyr Asn Pro Asn 305 \$310\$
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- Tyr Ala Arg Ser Tyr Cys Phe Ser Ala Asn Tyr Ile Tyr His Leu Phe 420 425 430
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Ile Phe Leu Ser Ser Met Cys Pro Ile Asn Val Ser Ala Ser Thr Leu 35 40 45

Tyr Gly Ile Met Phe Asp Ala Gly Ser Thr Gly Thr Arg Ile His Val $_{50}$

- Tyr Thr Phe Val Gln Lys Met Pro Gly Gln Leu Pro Ile Leu Glu Gly 65 70 75 80
- Glu Val Phe Asp Ser Val Lys Pro Gly Leu Ser Ala Phe Val Asp Gln 85 90 95
- Pro Lys Gln Gly Ala Glu Thr Val Gln Gly Leu Leu Glu Val Ala Lys
- Asp Ser Ile Pro Arg Ser His Trp Lys Lys Thr Pro Val Val Leu Lys 115 120 125
- Ala Thr Ala Gly Leu Arg Leu Leu Pro Glu His Lys Ala Lys Ala Leu 130 135 140
- Leu Phe Glu Val Lys Glu Ile Phe Arg Lys Ser Pro Phe Leu Val Pro 145 150 155 160
- Lys Gly Ser Val Ser Ile Met Asp Gly Ser Asp Glu Gly Ile Leu Ala 165 170 175
- Trp Val Thr Val Asn Phe Leu Thr Gly Gln Leu His Gly His Arg Gln
- Glu Thr Val Gly Thr Leu Asp Leu Gly Gly Ala Ser Thr Gln Ile Thr
- Phe Leu Pro Gln Phe Glu Lys Thr Leu Glu Gln Thr Pro Arg Gly Tyr 210 215 220
- Leu Thr Ser Phe Glu Met Phe Asn Ser Thr Tyr Lys Leu Tyr Thr His 225 230 235 240
- Ser Tyr Leu Gly Phe Gly Leu Lys Ala Ala Arg Leu Ala Thr Leu Gly 245 250 255
- Ala Leu Glu Thr Glu Gly Thr Asp Gly His Thr Phe Arg Ser Ala Cys 260 265 270
- Leu Pro Arg Trp Leu Glu Ala Glu Trp Ile Phe Gly Gly Val Lys Tyr 275 280 285
- Gln Tyr Gly Gly Asn Gln Glu Gly Glu Val Gly Phe Glu Pro Cys Tyr 290 295 300
- Ala Glu Val Leu Arg Val Val Arg Gly Lys Leu His Gln Pro Glu Glu 305 310 315
- Val Gln Arg Gly Ser Phe Tyr Ala Phe Ser Tyr Tyr Tyr Asp Arg Ala 325 330 335
- Val Asp Thr Asp Met Ile Asp Tyr Glu Lys Gly Gly Ile Leu Lys Val 340 345 350
- Glu Asp Phe Glu Arg Lys Ala Arg Glu Val Cys Asp Asn Leu Glu Asn 355 360 365

Phe Thr Ser Gly Ser Pro Phe Leu Cys Met Asp Leu Ser Tyr Ile Thr 370 380

Ala Leu Leu Lys Asp Gly Phe Gly Phe Ala Asp Ser Thr Val Leu Gln 385 390 395 400

Leu Thr Lys Lys Val Asn Asn Ile Glu Thr Gly Trp Ala Leu Gly Ala 405 410 415

Thr Phe His Leu Leu Gln Ser Leu Gly Ile Ser His
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<220>

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aagggaggge ctgaaggace tecacaggag tgtgagcage actgetteag caacaaagee 180

teaggteeac atcttgggaa gaat atg gcc act tee tgg ggg get gte tte 231
Met Ala Thr Ser Trp Gly Ala Val Phe

atg ctg atc ata gcc tgc gtt ggc agc act gtc ttc tac aga gaa cag 279 Met Leu Ile Ile Ala Cys Val Gly Ser Thr Val Phe Tyr Arg Glu Gln 10 15 20 25

cag acc tgg ttt gaa ggt gtc ttc ttg tct tcc atg tgc ccc att aat 327 Gln Thr Trp Phe Glu Gly Val Phe Leu Ser Ser Met Cys Pro Ile Asn 30 35 40

gtc agt gcc ggc acc ttt tat gga att atg ttt gat gcg ggc agc act 375 Val Ser Ala Gly Thr Phe Tyr Gly Ile Met Phe Asp Ala Gly Ser Thr 45 50

gga gct cgg att cat gtt tac act ttt gtg cag aaa aca gca gga cag 423 Gly Ala Arg Ile His Val Tyr Thr Phe Val Gln Lys Thr Ala Gly Gln 65 70

ctc ccc ttt ctg gaa ggt gaa att ttt gat tct gtg aag ccg gga ctt 471 Leu Pro Phe Leu Glu Gly Gly Ile Phe Asp Ser Val Lys Pro Gly Leu 75 80 85

tot got tit gig gat cag ccc aaa cag ggt gct gag act gic cag gag 519 Ser Ala Phe Val Asp Gln Pro Lys Gln Gly Ala Glu Thr Val Gln Glu 90 95 100

ctc Leu	ttg Leu	gag Glu	gtg Val	gcc Ala 110	Lys	gac Asp	tcg Ser	atc Ile	Pro	aga Arg	agc Ser	cac His	tgg T rp	gaa Glu 120	agg Arg	567
acc Thr	ccg Pro	gtg Val	gtt Val 125	ctg Leu	aaa Lys	gca Ala	acg Thr	gcc Ala 130	gga Gly	ctc Leu	cgt Arg	ttg Leu	ctg Leu 135	cct Pro	gag Glu	615
cag Gln	aaa Lys	gcc Ala 140	cag Gln	gct Ala	ctg Leu	ctc Leu	ttg Leu 145	gag Glu	gta Val	gag Glu	gag Glu	atc Ile 150	ttc Phe	aag Lys	aat Asn	663
tca Ser	cct Pro 155	ttc Phe	ctg Leu	gtc Val	cca Pro	gat Asp 160	ggc	agc Ser	gtt Val	agc Ser	atc Ile 165	atg Met	gat Asp	ggg ggg	tcc Ser	711
tat Tyr 170	gaa Glu	ggc Gly	ata Ile	cta Leu	gcc Ala 175	tgg Trp	gtt Val	acc Thr	gtg Val	aac Asn 180	ttt Phe	cta Leu	aca Thr	ggt Gly	cag Gln 185	759
ctg Leu	cat His	ggt Gly	cgt Arg	ggc Gly 190	cag Gln	gag Glu	act Thr	gtg Val	999 Gly 195	acc Thr	ctt Leu	gac Asp	ctg Leu	ggg Gly 200	ggt Gly	807
gcc Ala	tcc Ser	acc Thr	caa Gln 205	atc Ile	acg Thr	ttt Phe	cta Leu	ccc Pro 210	cag Gln	ttt Phe	gag Glu	aaa Lys	acc Thr 215	ctg Le u	gaa Glu	855
caa Gln	aca Thr	cct Pro 220	agg Arg	ggc Gly	tac Tyr	ctc Leu	act Thr 225	tcc Ser	ttt Phe	gag Glu	atg Met	ttt Phe 230	aac Asn	agc Ser	act Thr	903
ttt Phe	aag Lys 235	ctc Leu	tat Tyr	aca Thr	cat His	agt Ser 240	tac Tyr	ttg Leu	gga Gly	ttt Phe	gga Gly 245	ctg Leu	aaa Lys	gct Ala	gca Ala	951
aga Arg 250	ctg Leu	gca Ala	act Thr	ctg Leu	gga Gly 255	gcc Ala	ctg Leu	gaa Glu	gca Ala	aaa Lys 260	G1y 999	act Thr	gat Asp	gga Gly	cat His 265	999
acg Thr	ttt Phe	cga Arg	agt Ser	gcc Ala 270	tgt Cys	tta Leu	cca Pro	aga Arg	tgg Trp 275	ttg Leu	gaa Glu	gca Ala	gag Glu	tgg Trp 280	atc Ile	1047
ttt Phe	ggg ggg	ggt Gly	gtg Val 285	aaa Lys	tac Tyr	cag Gln	tat Tyr	ggt Gly 290	ggt Gly	aac Asn	caa Gln	gaa Glu	999 Gly 295	gag Glu	atg Met	1095
ggc Gly	ttt Phe	gaa Glu 300	ccc Pro	tgc Cys	tat Tyr	gcg Ala	gaa Glu 305	gtg Val	ctg Leu	agg Arg	gta Val	gta Val 310	cag Gln	ggg Gly	aaa Lys	1143
ctt Leu	cac His 3 15	cag Gln	cca Pro	gaa Glu	gaa Glu	gtc Val 320	cga Arg	gga Gly	agc Ser	gcc Ala	ttc Phe 325	tac Tyr	gct Ala	ttc Phe	tct Ser	1191

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														gaa Glu 360		1287
														tgc Cys		1335
														ttt Phe		1383
														aga Arg		1431
														ggg Gly		1479
														gaa Glu 440		1527
														ctc Leu		1575
				ccc Pro				taad	etggt	tt t	ataa	agga	39 ga	agggg	gtttt	1629
taga	atgag	gtc t	tgct	tctt	ga go	ctag	gtgat	tte	gggct	tca	atga	attt	gca (eatct	aatgt	1689
gaat	agct	cc 1	aaco	cact	g gt	gggt	gcat	gg	tgg	cacc	agad	tgta	aaa 1	ctti	tggga	1749
ttct	ttgt	ac a	agagt	tcct	gc aa	agga	aaaa	aga	agaaa	aagg	ttt	ggaa	ete (catgo	ctagat	1809
tgcg	gagtt	ca e	gaga	caggi	c c	tggg	ggac	aaa	agaad	caat	ctc	gttt	caa o	cccti	ggatg	1869
cct	catte	get 1	ttgaa	atgga	at to	catti	ttg	tta	ataa	gctg	att	tact	gaa a	atcc	cataac	1929
ccat	caat	tgc 1	tgtta	aatti	tt ti	tcti	ccta	a cc	ctta	tac	att	ccct	acc	ctaa	aagcct	1989
9999	ggaaa	ata (cctg	gttt	tg c	ttee	catc	ata	aatt	gaga	aag	aggg	ggg (aaaa	gatact	2049
gtat	ttaga	aat	ttgt	gtgai	te e	gtg	gcaca	ata	agat	caac	caa	ccca	ttt .	aaag	cttaaa	2109
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Met Ala Thr Ser Trp Gly Ala Val Phe Met Leu lie ile Ala Cys Val 1 5 10 15

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Phe Leu Ser Ser Met Cys Pro Ile Asn Val Ser Ala Gly Thr Phe Tyr \$35\$ \$40\$ \$45\$

Gly Ile Met Phe Asp Ala Gly Ser Thr Gly Ala Arg Ile His Val Tyr 50 60

Thr Phe Val Gln Lys Thr Ala Gly Gln Leu Pro Phe Leu Glu Gly Glu 65 70 75 80

Ile Phe Asp Ser Val Lys Pro Gly Leu Ser Ala Phe Val Asp Gln Pro 85 90 95

Lys Gln Gly Ala Glu Thr Val Gln Glu Leu Leu Glu Val Ala Lys Asp $100 \hspace{1.5cm} 105 \hspace{1.5cm} 110 \hspace{1.5cm}$

Ser Ile Pro Arg Ser His Trp Glu Arg Thr Pro Val Val Leu Lys Ala 115 120 125

Thr Ala Gly Leu Arg Leu Leu Pro Glu Gln Lys Ala Gln Ala Leu Leu 130 135 140

Leu Glu Val Glu Glu Ile Phe Lys Asn Ser Pro Phe Leu Val Pro Asp 145 150 155 160

Gly Ser Val Ser Ile Met Asp Gly Ser Tyr Glu Gly Ile Leu Ala Trp 165 170 175

Val Thr Val Asn Phe Leu Thr Gly Gln Leu His Gly Arg Gly Gln Glu 180 185 190

Thr Val Gly Thr Leu Asp Leu Gly Gly Ala Ser Thr Gln Ile Thr Phe 195 200

Leu Pro Gln Phe Glu Lys Thr Leu Glu Gln Thr Pro Arg Gly Tyr Leu 210 215220

Thr Ser Phe Glu Met Phe Asn Ser Thr Phe Lys Leu Tyr Thr His Ser 225 230 235 240

Leu Glu Ala Lys Gly Thr Asp Gly His Thr Phe Arg Ser Ala Cys Leu 260 265 270

Pro Arg Trp Leu Glu Ala Glu Trp Ile Phe Gly Gly Val Lys Tyr Gln 275 280 285

Tyr Gly Gly Asn Gln Glu Gly Glu Met Gly Phe Glu Pro Cys Tyr Ala 290 295 300 Glu Val Leu Arg Val Val Gln Gly Lys Leu His Gln Pro Glu Glu Val 305 315 315 326

Arg Gly Ser Ala Phe Tyr Ala Phe Ser Tyr Tyr Tyr Asp Arg Ala Ala

Asp Thr His Leu Ile Asp Tyr Glu Lys Gly Gly Val Leu Lys Val Glu 340 345 350

Asp Phe Glu Arg Lys Ala Arg Glu Val Cys Asp Asn Leu Gly Ser Phe 355 360 365

Ser Ser Gly Ser Pro Phe Leu Cys Met Asp Leu Thr Tyr Ile Thr Ala 370 380

Leu Leu Lys Asp Gly Leu Gly Phe Ala Glu Arg His Pro Leu Thr Ala 385 390 395 400

His Lys Glu Ser Glu Gln His Arg Asp Trp Leu Gly Leu Gly Gly His

Leu Ser Pro Ala Pro Val Ser Gly His His Gln Leu Arg Pro Ser Ser 420 425 430

Thr Ser Glu Ala Cys Ile Ser Glu Pro Val Phe Ser Gln Glu Gly Val 435 \$440\$

Asp Ser Glu Thr Phe Ser Asp Leu Ser Gly Lys Ala Trp Pro Glu Thr 450 $\,$ 455 $\,$ 460

Arg 465

<210> 9 <211> 428 <212> PRT <213> Homo sapiens

<400> 9

Met Ala Thr Ser Trp Gly Thr Val Phe Phe Met Leu Val Val Ser Cys

1 10 15

Val Cys Ser Ala Val Ser His Arg Asn Gln Gln Thr Trp Phe Glu Gly
20 25 30

Ile Phe Leu Ser Ser Met Cys Pro Ile Asn Val Ser Ala Ser Thr Leu

Tyr Gly Ile Met Phe Asp Ala Gly Ser Thr Gly Thr Arg Ile His Val $_{50}^{\rm co}$

Tyr Thr Phe Val Gln Lys Met Pro Gly Gln Leu Pro Ile Leu Glu Gly 65 70 75 80

- Glu Val Phe Asp Ser Val Lys Pro Gly Leu Ser Ala Phe Val Asp Gln 85 90 95
- Pro Lys Gln Gly Ala Glu Thr Val Gln Gly Leu Leu Glu Val Ala Lys 100 105 110
- Asp Ser Ile Pro Arg Ser His Trp Lys Lys Thr Pro Val Val Leu Lys 115 120 125
- Ala Thr Ala Gly Leu Arg Leu Leu Pro Glu His Lys Ala Lys Ala Leu
 130 135 140
- Leu Phe Glu Val Lys Glu Ile Phe Arg Lys Ser Pro Phe Leu Val Pro 145 150 155 160
- Lys Gly Ser Val Ser Ile Met Asp Gly Ser Asp Glu Gly Ile Leu Ala
- Trp Val Thr Val Asn Phe Leu Thr Gly Gln Leu His Gly His Arg Gln
- Glu Thr Val Gly Thr Leu Asp Leu Gly Gly Ala Ser Thr Gln Ile Thr
- Phe Leu Pro Gln Phe Glu Lys Thr Leu Glu Gln Thr Pro Arg Gly Tyr
- Leu Thr Ser Phe Glu Met Phe Asn Ser Thr Tyr Lys Leu Tyr Thr His 225 230 235 240
- Ser Tyr Leu Gly Phe Gly Leu Lys Ala Ala Arg Leu Ala Thr Leu Gly 245 250 255
- Ala Leu Glu Thr Glu Gly Thr Asp Gly His Thr Phe Arg Ser Ala Cys 260 265 270
- Leu Pro Arg Trp Leu Glu Ala Glu Trp Ile Phe Gly Gly Val Lys Tyr
 275 280 285
- Gln Tyr Gly Gly Asn Gln Glu Gly Glu Val Gly Phe Glu Pro Cys Tyr 290 295 300
- Ala Glu Val Leu Arg Val Val Arg Gly Lys Leu His Gln Pro Glu Glu 305 310 315 320
- Val Gln Arg Gly Ser Phe Tyr Ala Phe Ser Tyr Tyr Tyr Asp Arg Ala 325 330 335
- Val Asp Thr Asp Met Ile Asp Tyr Glu Lys Gly Gly Ile Leu Lys Val 340 345 350
- Glu Asp Phe Glu Arg Lys Ala Arg Glu Val Cys Asp Asn Leu Glu Asn 355 360 365
- Phe Thr Ser Gly Ser Pro Phe Leu Cys Met Asp Leu Ser Tyr Ile Thr 370 375 380

Ala Leu Leu Lys Asp Gly Phe Gly Phe Ala Asp Ser Thr Val Leu Gln 385 390 395

Leu Thr Lys Lys Val Asn Asn Ile Glu Thr Gly Trp Ala Leu Gly Ala
405 410 415

Thr Phe His Leu Leu Gln Ser Leu Gly Ile Ser His
420 425

<210> 10

<211> 455

<212> PRT

<213> P. sativum

<400> 10

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Ala Ile Thr Ser Ser Gln Tyr Leu Gly Asn Asn Leu Leu Thr Ser Arg

Lys Ile Phe Leu Lys Gln Glu Glu Ile Ser Ser Tyr Ala Val Val Phe $35 \hspace{1cm} 40 \hspace{1cm} 45$

Asp Ala Gly Ser Thr Gly Ser Arg Ile His Val Tyr His Phe Asn Gln 50 55 60

Asn Leu Asp Leu Leu His Ile Gly Lys Gly Val Glu Tyr Tyr Asn Lys 65 70 75 80

Ile Thr Pro Gly Leu Ser Ser Tyr Ala Asn Asn Pro Glu Gln Ala Ala 85 90 95

Lys Ser Leu Ile Pro Leu Leu Glu Gln Ala Glu Asp Val Val Pro Asp 100 105 110

Asp Leu Gln Pro Lys Thr Pro Val Arg Leu Gly Ala Thr Ala Gly Leu 115 120 125

Arg Leu Leu Asn Gly Asp Ala Ser Glu Lys Ile Leu Gln Ser Val Arg 130 135 140

Asp Met Leu Ser Asn Arg Ser Thr Phe Asn Val Gln Pro Asp Ala Val 145 150 155 160

Ser Ile Ile Asp Gly Thr Gln Glu Gly Ser Tyr Leu Trp Val Thr Val 165 170 175

Asn Tyr Ala Leu Gly Asn Leu Gly Lys Lys Tyr Thr Lys Thr Val Gly 180 185 190

Val Ile Asp Leu Gly Gly Gly Ser Val Gln Met Ala Tyr Ala Val Ser 195 200 205

Lys Lys Thr Ala Lys Asn Ala Pro Lys Val Ala Asp Gly Asp Asp Pro 210 215 220

Tyr Ile Lys Lys Val Val Leu Lys Gly Ile Pro Tyr Asp Leu Tyr Val 225 230 235 240

His Ser Tyr Leu His Phe Gly Arg Glu Ala Ser Arg Ala Glu Ile Leu 245 250 255

Gly Ile Tyr Thr Tyr Ser Gly Glu Glu Phe Lys Ala Thr Ala Tyr Thr $275 \hspace{1cm} 280 \hspace{1cm} 285$

Ser Gly Ala Asn Phe Asn Lys Cys Lys Asn Thr Ile Arg Lys Ala Leu 290 295 300

Lys Leu Asn Tyr Pro Cys Pro Tyr Gln Asn Cys Thr Phe Gly Gly Ile 305 310 315 320

Trp Asn Gly Gly Gly Gly Asn Gly Gln Lys Asn Leu Phe Ala Ser Ser $325 \hspace{1.5cm} 330 \hspace{1.5cm} 335 \hspace{1.5cm}$

Ser Phe Phe Tyr Leu Pro Glu Asp Thr Gly Met Val Asp Ala Ser Thr 340 345 350

Pro Asn Phe Ile Leu Arg Pro Val Asp Ile Glu Thr Lys Ala Lys Glu 355 360 365

Ala Cys Ala Leu Asn Phe Glu Asp Ala Lys Ser Thr Tyr Pro Phe Leu 370 380

Asp Lys Lys Asn Val Ala Ser Tyr Val Cys Met Asp Leu Ile Tyr Gln 385 390395395

Thr Ser Gly Lys Glu Ile Glu Tyr Gln Asp Ala Ile Val Glu Ala Ala 420 \$425\$

Trp Pro Leu Gly Asn Ala Val Glu Ala Ile Ser Ala Leu Pro Lys Phe $435 \ \ \,$ 440 $\ \ \,$

Glu Arg Leu Met Tyr Phe Val 450 455

<210> 11

<211> 454

<212> PRT

<213> Solanum tuberosum

<400> 11

Met Leu Asn Gln Asn Ser His Phe Ile Phe Ile Ile Leu Ala Ile Phe 1 10 15

Leu Val Leu Pro Leu Ser Leu Leu Ser Lys Asn Val Asn Ala Gln Ile 20 25 30 Pro Leu Arg Arg His Leu Leu Ser His Glu Ser Glu His Tyr Ala Val

Ile Phe Asp Ala Gly Ser Thr Gly Ser Arg Val His Val Phe Arg Phe

Asp Glu Lys Leu Gly Leu Leu Pro Ile Gly Asn Asn Ile Glu Tyr Phe 65 70 75 80

Met Ala Thr Glu Pro Gly Leu Ser Ser Tyr Ala Glu Asp Pro Lys Ala 85 90 95

Ala Ala As
n Ser Leu Glu Pro Leu Leu Asp Gly Ala Glu Gly Val Val 100 105 110

Pro Gln Glu Leu Gln Ser Glu Thr Pro Leu Glu Leu Gly Ala Thr Ala 115 120 125

Gly Leu Arg Met Leu Lys Gly Asp Ala Ala Glu Lys Ile Leu Gln Ala 130 135 140

Val Arg Asn Leu Val Lys Asn Gln Ser Thr Phe His Ser Lys Asp Gln 145 150 155 160

Trp Val Thr Ile Leu Asp Gly Thr Gln Glu Gly Ser Tyr Met Trp Ala 165 170 175

Thr Ala Thr Ile Asp Leu Gly Gly Gly Ser Val Gln Met Ala Tyr Ala 195 200 205

Ile Ser Asn Glu Gln Phe Ala Lys Ala Pro Gln Asn Glu Asp Gly Glu 210 215 220

Pro Tyr Val Gln Gln Lys His Leu Met Ser Lys Asp Tyr Asn Leu Tyr 225 $$ 230 $$ 235 $$ 240

Val His Ser Tyr Leu Asn Tyr Gly Gln Leu Ala Gly Arg Ala Glu Ile 245 250 255

Phe Lys Ala Ser Arg Asn Glu Ser Asn Pro Cys Ala Leu Glu Gly Cys

Asp Gly Tyr Tyr Ser Tyr Gly Gly Val Asp Tyr Lys Val Lys Ala Pro

Lys Lys Gly Ser Ser Trp Lys Arg Cys Arg Arg Leu Thr Arg His Ala 290 295 300

Leu Lys Ile Asn Ala Lys Cys Asn Ile Glu Glu Cys Thr Phe Asn Gly 305 310 315 320

Val Trp Asn Gly Gly Gly Gly Asp Gly Gln Lys Asn Ile His Ala Ser 325 330 335

Ser Phe Phe Tyr Asp Ile Gly Ala Gln Val Gly Ile Val Asp Thr Lys \$340\$

Phe Pro Ser Ala Leu Ala Lys Pro Ile Gln Tyr Leu Asn Ala Ala Lys 355 360 365

Val Ala Cys Gln Thr Asn Val Ala Asp Ile Lys Ser Ile Phe Pro Lys 370 375 380

Thr Gln Asp Arg Asn Ile Pro Tyr Leu Cys Met Asp Leu Ile Tyr Glu 385 390 395 400

Tyr Thr Leu Leu Val Asp Gly Phe Gly Leu Asn Pro His Lys Glu Ile 405 410 415

Thr Val Ile His Asp Val Gln Tyr Lys Asn Tyr Leu Val Gly Ala Ala 420 \$425\$

Ile Arg Val Ala Ser Ser 450

<210> 12

<211> 473

<212> PRT

<213> Saccharomyces cerevisiae

<400> 12

Tyr Leu Gln Asp Ser Lys Thr Glu Gln Asn Tyr Pro Glu Leu Ala Asp 20 25 30

Ala Val Lys Ser Gln Thr Ser Gln Thr Cys Ser Glu Glu His Lys Tyr $35 \hspace{1cm} 40 \hspace{1cm} 45 \hspace{1cm}$

Val Ile Met Ile Asp Ala Gly Ser Thr Gly Ser Arg Val His Ile Tyr 50 55 60

Lys Phe Asp Val Cys Thr Ser Pro Pro Thr Leu Leu Asp Glu Lys Phe 65 70 75 80

Asp Met Leu Glu Pro Gly Leu Ser Ser Phe Asp Thr Asp Ser Val Gly

Ala Ala Asn Ser Leu Asp Pro Leu Leu Lys Val Ala Met Asn Tyr Val

Pro Ile Lys Ala Arg Ser Cys Thr Pro Val Ala Val Lys Ala Thr Ala 115 120 125

Gly Leu Arg Leu Leu Gly Asp Ala Lys Ser Ser Lys Ile Leu Ser Ala 130 \$135\$

- Val Arg Asp His Leu Glu Lys Asp Tyr Pro Phe Pro Val Val Glu Gly 145 150 155 160
- Asp Gly Val Ser Ile Met Gly Gly Asp Glu Glu Gly Val Phe Ala Trp 165 170 175
- Ile Thr Thr Asn Tyr Leu Leu Gly Asn Ile Gly Ala Asn Gly Pro Lys 180 185 190
- Leu Pro Thr Ala Ala Val Phe Asp Leu Gly Gly Gly Ser Thr Gln Ile 195 200 205
- Val Glu Glu Pro Thr Phe Pro Ile Asn Glu Lys Met Val Asp Gly Glu 210 215 220
- His Lys Phe Asp Leu Lys Phe Gly Asp Glu Asn Tyr Thr Leu Tyr Gln 225 230 235 240
- Phe Ser His Leu Gly Tyr Gly Leu Lys Glu Gly Arg Asn Lys Val Asn 245 250 255
- Ser Val Leu Val Glu Asn Ala Leu Lys Asp Lys Ile Leu Lys Gly Cys
- Asn Thr Lys Thr His Cys Leu Ser Ser Pro Cys Leu Pro Pro Lys Val
- Asn Ala Thr Asn Glu Lys Val Thr Leu Glu Ser Lys Glu Thr Tyr Thr 290 295 300
- Ile Asp Phe Ile Gly Pro Asp Glu Pro Ser Gly Ala Gln Cys Arg Phe 305 \$310\$
- Leu Thr Asp Glu Ile Leu Asn Lys Asp Ala Gln Cys Gln Ser Pro Pro 325 330 335
- Cys Ser Phe Asn Gly Val His Gln Pro Ser Leu Val Arg Thr Phe Lys 340 345 350
- Glu Ser Asn Asp Ile Tyr Ile Phe Ser Tyr Phe Tyr Asp Arg Thr Thr 355 \$360\$
- Arg Pro Leu Gly Met Pro Leu Ser Phe Thr Leu Asn Glu Leu Asn Asp 370 375 380
- Leu Ala Arg Ile Val Cys Lys Gly Glu Glu Thr Trp Asn Ser Val Phe 385 \$390 , \$395
- Ser Gly Ile Ala Gly Ser Leu Asp Glu Leu Glu Ser Asp Ser His Phe 405 410 415
- Cys Leu Asp Leu Ser Phe Gln Val Ser Leu Leu His Thr Gly Tyr Asp 420 425 430
- Ile Pro Leu Gln Arg Glu Leu Arg Thr Gly Lys Lys Ile Ala Asn Lys 435 440 445

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Asn Trp Lys Cys Lys Ile Gln Ser Ala
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<211> 153
<212> PRT
<213> Homo sapiens
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Ile Tyr Lys Trp Pro Ala Glu Lys Glu Asn Asp Thr Gly Val Val His
Gln Val Glu Glu Cys Arg Val Lys Gly Pro Gly Ile Ser Lys Phe Val
Gln Lys Val Asn Glu Ile Gly Ile Tyr Leu Thr Asp Cys Met Glu Arg
Ala Arg Glu Val Ile Pro Arg Ser Gln His Gln Glu Thr Pro Val Tyr
Leu Gly Ala Thr Ala Gly Met Arg Leu Leu Arg Met Glu Ser Glu Glu
Leu Ala Asp Arg Val Leu Asp Val Val Glu Arg Ser Leu Ser Asn Tyr
Pro Phe Asp Phe Gln Gly Ala Arg Ile Ile Thr Gly Gln Glu Gly
        115
Ala Tyr Gly Trp Ile Thr Ile Asn Tyr Leu Leu Gly Lys Phe Ser Gln
                        135
Lys Thr Arg Trp Phe Ser Ile Val Pro
                    150
145
<210> 14
<211> 154
<212> PRT
<213> Rattus norvegicus
<400> 14
Val Lys Tyr Gly Ile Val Leu Asp Ala Gly Ser Ser His Thr Asn Leu
Tyr Ile Tyr Lys Trp Pro Ala Glu Lys Glu Asn Asp Thr Gly Val Val
Gln Leu Leu Glu Glu Cys Gln Val Lys Gly Pro Gly Ile Ser Lys Tyr
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Ala Gln Lys Thr Asp Glu Ile Ala Ala Tyr Leu Ala Glu Cys Met Lys 50 60

Met Ser Thr Glu Arg Ile Pro Ala Ser Lys Gln His Gln Thr Pro Val 65 70 75 80

Tyr Leu Gly Ala Thr Ala Gly Met Arg Leu Leu Arg Met Glu Ser Lys 90 95

Gln Ser Ala Asp Glu Val Leu Ala Ala Val Ser Arg Ser Leu Lys Ser

Tyr Pro Phe Asp Phe Gln Gly Ala Lys Ile Ile Thr Gly Gln Glu Glu 115 120 125

Gly Ala Tyr Gly Trp Ile Thr Ile Asn Tyr Leu Leu Gly Arg Phe Thr 130 $$135\$

Gln Glu Gln Ser Trp Leu Asn Phe Ile Ser 145

<210> 15

<211> 153 <212> PRT

<213> Homo sapiens

<400> 15

Lys Tyr Gly Ile Val Leu Asp Ala Gly Ser Ser His Thr Ser Met Phe 1 5 10 15

Ile Tyr Lys Trp Pro Ala Asp Lys Glu Asn Asp Thr Gly Ile Val Gly 20 25 30

Gln His Ser Ser Cys Asp Val Pro Gly Gly Gly Ile Ser Ser Tyr Ala 35 4045

Asp Asn Pro Ser Gly Ala Ser Gln Ser Leu Val Gly Cys Leu Glu Gln 50 55 60

Ala Leu Gln Asp Val Pro Lys Glu Arg His Ala Gly Thr Pro Leu Tyr 65 70 75 80

Leu Gly Ala Thr Ala Gly Met Arg Leu Leu Asn Leu Thr Asn Pro Glu 85 90 95

Ala Ser Thr Ser Val Leu Met Ala Val Thr His Thr Leu Thr Gln Tyr 100 $$105\$

Pro Phe Asp Phe Arg Gly Ala Arg Ile Leu Ser Gly Glu Glu Gly 115 120 125

Tyr Gly Trp Val Gly Arg Trp Phe Arg

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<211> 150
<212> PRT
<213> Gallus gallus
<400> 16
Phe Lys Tyr Gly Ile Val Leu Asp Ala Gly Ser Ser His Thr Ala Val
Phe Ile Tyr Lys Trp Pro Ala Asp Lys Glu Asn Asp Thr Gly Val Val
Ser Glu His Ser Met Cys Asp Val Glu Gly Pro Gly Ile Ser Ser Tyr
Ser Ser Lys Pro Pro Ala Ala Gly Lys Ser Leu Glu His Cys Leu Ser
Gln Ala Met Arg Asp Val Pro Lys Glu Lys His Ala Asp Thr Pro Leu
Tyr Leu Gly Ala Thr Ala Gly Met Arg Leu Leu Thr Ile Ala Asp Pro
Pro Ser Gln Thr Cys Leu Ser Ala Val Met Ala Thr Leu Lys Ser Tyr
Pro Phe Asp Phe Gly Gly Ala Lys Ile Leu Ser Gly Glu Glu Gly
Val Phe Gly Trp Ile Thr Ala Asn Tyr Leu Leu Glu Asn Phe Ile Lys
Arg Gly Trp Leu Gly Glu
<210> 17
<211> 148
<212> PRT
<213> Caenorhabditis elegans
<400> 17
Ile Lys Tyr Gly Val Ile Cys Asp Ala Gly Ser Ser Gly Thr Arg Leu
Phe Val Tyr Thr Leu Lys Pro Leu Ser Gly Gly Leu Thr Asn Ile Asp
Thr Leu Ile His Glu Ser Glu Pro Val Val Lys Lys Val Thr Pro Gly
Leu Ser Ser Phe Gly Asp Lys Pro Glu Gln Val Val Glu Tyr Leu Thr
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Pro Leu Leu Arg Phe Ala Glu Glu His Ile Pro Tyr Glu Gln Leu Gly

70

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Glu Thr Asp Leu Leu Ile Phe Ala Thr Ala Gly Met Arg Leu Leu Pro
Glu Ala Gln Lys Asp Ala Ile Ile Lys Asn Leu Gln Asn Gly Leu Lys
            100
                                105
Ser Val Thr Ala Leu Arg Val Ser Asp Ser Asn Ile Arg Ile Ile Asp
Gly Ala Trp Glu Gly Ile Tyr Ser Trp Ile Ala Val Asn Tyr Ile Leu
Gly Arg Phe Asp
145
<210> 18
<211> 10
<212> RNA
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aagaauaugg
                                                                   10
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